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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/067,122B

DATE: 11/26/2002 TIME: 13:56:38

Input Set : A:\iu-0008.ST25.txt

Output Set: N:\CRF4\11262002\J067122B.raw

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3 <110> APPLICANT: Kwon, Byoung S
 5 <120> TITLE OF INVENTION: Antibody for 4-1BB
 7 <130> FILE REFERENCE: 740.009US2 (IU-0008)
 9 <140> CURRENT APPLICATION NUMBER: US 10/067,122B
10 <141> CURRENT FILING DATE: 2002-02-04
12 <150> PRIOR APPLICATION NUMBER: US 08/012,269
13 <151> PRIOR FILING DATE: 1993-02-01
15 <150> PRIOR APPLICATION NUMBER: US 07/922,996
16 <151> PRIOR FILING DATE: 1992-07-30
18 <150> PRIOR APPLICATION NUMBER: US 07/267,577
19 <151> PRIOR FILING DATE: 1988-11-07
21 <160> NUMBER OF SEQ ID NOS: 13
23 <170> SOFTWARE: PatentIn version 3.1
25 <210> SEO ID NO: 1
26 <211> LENGTH: 2350
27 <212> TYPE: DNA
28 <213> ORGANISM: Mus musculus
30 <220> FEATURE:
31 <221> NAME/KEY: misc feature
32 <222> LOCATION: (1253)..(1255)
33 <223> OTHER INFORMATION: n=a, c, g, or t
36 <400> SEQUENCE: 1
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39 ctacaccagg aaaaggacac attcgacaac aggaaaggag cctgtcacag aaaaccacag
41 tgtcctgtgc atgtgacatt tcgccatggg aaacaactgt tacaacgtgg tggtcattgt
                                                                         180
43 gctgctgcta gtgggctgtg agaaggtggg agccgtgcag aactcctgtg ataactgtca
                                                                         240
45 gcctggtact ttctgcagaa aatacaatcc agtctgcaag agctgccctc caagtacctt
                                                                         300
47 ctccagcata ggtggacagc cgaactgtaa catctgcaga gtgtgtgcag gctatttcag
                                                                         360
49 gttcaagaag ttttgctcct ctacccacaa cgcggagtgt gagtgcattg aaggattcca
                                                                         420
51 ttgcttgggg ccacagtgca ccagatgtga aaaggactgc aggcctggcc aggagctaac
                                                                         480
                                                                         540
53 gaagcagggt tgcaaaacct gtagettggg aacatttaat gaccagaacg gtactggcgt
                                                                         600
55 ctgtcgaccc tggacgaact gctctctaga cggaaggtct gtgcttaaga ccgggaccac
57 ggagaaggac gtggtgtgtg gaccccctgt ggtgagcttc tctcccagta ccaccatttc
                                                                         660
                                                                         720
59 tgtgactcca gagggaggac caggagggca ctccttgcag gtccttacct tgttcctggc
61 gctgacatcg gctttgctgc tggccctgat cttcattact ctcctgttct ctgtgctcaa
                                                                         780
                                                                         840
63 atqqatcagg aaaaaattcc cccacatatt caagcaacca tttaagaaga ccactggagc
                                                                         900
65 ageteaagag gaagatgett gtagetgeeg atgteeacag gaagaagaag gaggaggagg
                                                                         960
67 agqctatgag ctqtgatgta ctatcctagg agatgtgtgg gccgaaaccg agaagcacta
                                                                        1020
69 ggaccccacc atcctgtgga acagcacaag caaccccacc accctgttct tacacatcat
                                                                        1080
71 cctagatgat gtgtgggcgc gcacctcatc caagtctctt ctaacgctaa catatttgtc
73 tttacctttt ttaaatcttt ttttaaattt aaattttatg tgtgtgagtg ttttgcctgc
                                                                        1140
                                                                        1200
75 ctqtatqcac acqtqtqtqt qtqtqtqtqt qtqacactcc tqatqcctqa qgaqqtcaqa
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W--> 77 aqaqaaaggg ttggttccat aagaactgga gttatggatg gctgtgagcc ggnnngatag

1260

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79 qtcqqqacqq aqacctgtct tcttatttta acgtgactgt ataataaaaa aaaaatgata
                                                                        1320
81 tttcqqqaat tqtaqaqatt ctcctqacac ccttctaqtt aatqatctaa qaqqaattqt
                                                                        1380
83 tgatacgtag tatactgtat atgtgtatgt atatgtatat gtatatataa gactctttta
                                                                        1440
85 ctqtcaaaqt caacctaqaq tqtctqqtta ccaqqtcaat tttattqqac attttacqtc
                                                                        1500
87 acacacacac acacacacac acacacacqt ttatactacq tactqttatc qqtattctac
                                                                        1560
89 gtcatataat gggatagggt aaaaggaaac caaagagtga gtgatattat tgtggaggtg
                                                                        1620
91 acagactace cettetgggt acgtagggae agaceteett eggactgtet aaaacteece
                                                                        1680
93 ttagaagtct cgtcaagttc ccggacgaag aggacagagg agacacagtc cgaaaagtta
                                                                        1740
95 tttttccggc aaatcctttc cctgtttcgt gacactccac cccttgtgga cacttgagtg
                                                                        1800
97 teateettge geeggaaggt caggtggtae cegtetgtag gggcggggag acagageege
                                                                        1860
99 gggggagcta cgagaatcga ctcacagggc gccccgggct tcgcaaatga aactttttta
                                                                        1920
101 atctcacaag tttcgtccgg gctcggcgga cctatggcgt cgatccttat taccttatcc
                                                                         1980
103 tggcgccaag ataaaacaac caaaagcett gacteeggta etaattetee etgeeggeee
                                                                         2040
105 ccgtaagcat aacgcggcga tetecaettt aagaacetgg ccgcgttetg cetggteteg
                                                                         2100
107 ctttcgtaaa cggttcttac aaaagtaatt agttcttgct ttcagcctcc aagcttctgc
                                                                         2160
109 tagtctatgg cagcatcaag gctggtattt gctacggctg accgctacgc cgccgcaata
                                                                         2220
111 agggtactgg gcggcccgtc gaaggccctt tggtttcaga aacccaaggc cccctcata
                                                                         2280
113 ccaacgtttc gactttgatt cttgccggta cgtggtggtg ggtgccttag ctctttctcg
                                                                         2340
115 atagttagac
                                                                         2350
118 <210> SEQ ID NO: 2
119 <211> LENGTH: 256
120 <212> TYPE: PRT
121 <213> ORGANISM: Mus musculus
123 <400> SEQUENCE: 2
125 Met Gly Asn Asn Cys Tyr Asn Val Val Ile Val Leu Leu Val
126 1
                                         10
129 Gly Cys Glu Lys Val Gly Ala Val Gln Asn Ser Cys Asp Asn Cys Gln
                20
                                    25
133 Pro Gly Thr Phe Cys Arg Lys Tyr Asn Pro Val Cys Lys Ser Cys Pro
134
            35
                                40
137 Pro Ser Thr Phe Ser Ser Ile Gly Gly Gln Pro Asn Cys Asn Ile Cys
                            55
141 Arg Val Cys Ala Gly Tyr Phe Arg Phe Lys Lys Phe Cys Ser Ser Thr
142 65
                        70
                                            75
145 His Asn Ala Glu Cys Glu Cys Ile Glu Gly Phe His Cys Leu Gly Pro
                                        90
149 Gln Cys Thr Arg Cys Glu Lys Asp Cys Arg Pro Gly Gln Glu Leu Thr
150
                                    105
153 Lys Gln Gly Cys Lys Thr Cys Ser Leu Gly Thr Phe Asn Asp Gln Asn
154
            115
                                120
157 Gly Thr Gly Val Cys Arg Pro Trp Thr Asn Cys Ser Leu Asp Gly Arg
158
        130
                            135
                                                 140
161 Ser Val Leu Lys Thr Gly Thr Thr Glu Lys Asp Val Val Cys Gly Pro
162 145
                        150
                                            155
165 Pro Val Val Ser Phe Ser Pro Ser Thr Thr Ile Ser Val Thr Pro Glu
166
                    165
                                        170
169 Gly Gly Pro Gly Gly His Ser Leu Gln Val Leu Thr Leu Phe Leu Ala
                180
                                    185
173 Leu Thr Ser Ala Leu Leu Leu Ala Leu Ile Phe Ile Thr Leu Leu Phe
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```
174
                195
     177 Ser Val Leu Lys Trp Ile Arg Lys Lys Phe Pro His Ile Phe Lys Gln
                                 215
     181 Pro Phe Lys Lys Thr Thr Gly Ala Ala Gln Glu Glu Asp Ala Cys Ser
                             230
                                                 235
     185 Cys Arg Cys Pro Gln Glu Glu Glu Gly Gly Gly Gly Tyr Glu Leu
                                             250
     189 <210> SEQ ID NO: 3
     190 <211> LENGTH: 24
     191 <212> TYPE: PRT
     192 <213> ORGANISM: Mus musculus
     194 <400> SEQUENCE: 3
     196 Cys Arg Val Cys Ala Gly Tyr Phe Arg Phe Lys Lys Phe Cys Ser Ser
                                             10
     200 Thr His Asn Ala Glu Cys Glu Cys
     201
                    20
     204 <210> SEQ ID NO: 4
     205 <211> LENGTH: 22
     206 <212> TYPE: PRT
     207 <213> ORGANISM: Drosophila
     209 <400> SEQUENCE: 4
     211 Cys Pro Val Cys Phe Asp Tyr Val Ile Leu Gln Cys Ser Ser Gly His
     212 1
     215 Leu Val Cys Val Ser Cys
     216
                     20
     219 <210> SEQ ID NO: 5
     220 <211> LENGTH: 26
     221 <212> TYPE: PRT
     222 <213> ORGANISM: Dictyostelium
     224 <400> SEQUENCE: 5
     226 Cys Pro Ile Cys Phe Glu Phe Ile Tyr Lys Lys Gln Ile Tyr Gln Cys
                    5
     230 Lys Ser Gly His His Ala Cys Lys Glu Cys
     231
                    20
    234 <210> SEQ ID NO: 6
    235 <211> LENGTH: 6
     236 <212> TYPE: PRT
     237 <213> ORGANISM: Mus musculus
    239 <220> FEATURE:
    240 <221> NAME/KEY: MISC FEATURE
     241 <222> LOCATION: (5)..(5)
     242 <223> OTHER INFORMATION: X=any amino acid
     245 <400> SEQUENCE: 6
W--> 247 Val Gln Asn Ser Xaa Asp
     248 1
    251 <210> SEQ ID NO: 7
    252 <211> LENGTH: 12
    253 <212> TYPE: PRT
    254 <213> ORGANISM: Artificial Sequence
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Input Set : A:\iu-0008.ST25.txt Output Set: N:\CRF4\11262002\J067122B.raw 256 <220> FEATURE: 257 <223> OTHER INFORMATION: Synthetic 259 <400> SEQUENCE: 7 261 Cys Arg Pro Gly Gln Glu Leu Thr Lys Ser Gly Tyr 265 <210> SEQ ID NO: 8 266 <211> LENGTH: 24 267 <212> TYPE: PRT 268 <213> ORGANISM: Artificial Sequence 270 <220> FEATURE: 271 <223> OTHER INFORMATION: Synthetic 273 <220> FEATURE: 274 <221> NAME/KEY: MISC_FEATURE 275 <222> LOCATION: (2)..(3) 276 <223> OTHER INFORMATION: X=any amino acid 279 <220> FEATURE: 280 <221> NAME/KEY: MISC FEATURE 281 <222> LOCATION: (5)..(13) 282 <223> OTHER INFORMATION: X=any amino acid 285 <220> FEATURE: 286 <221> NAME/KEY: MISC FEATURE 287 <222> LOCATION: (15)..(17) 288 <223> OTHER INFORMATION: X=any amino acid 291 <220> FEATURE: 292 <221> NAME/KEY: MISC FEATURE 293 <222> LOCATION: (19)..(21) 294 <223> OTHER INFORMATION: X=any amino acid 297 <220> FEATURE: 298 <221> NAME/KEY: MISC FEATURE 299 <222> LOCATION: (23)..(23) 300 <223> OTHER INFORMATION: X=any amino acid 303 <400> SEQUENCE: 8 W--> 305 Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa 306 1 10 W--> 309 Xaa His Xaa Xaa Xaa Cys Xaa Cys 20 313 <210> SEO ID NO: 9 314 <211> LENGTH: 4 315 <212> TYPE: PRT 316 <213> ORGANISM: Artificial Sequence 318 <220> FEATURE: 319 <223> OTHER INFORMATION: Synthetic 321 <400> SEQUENCE: 9 323 Cys Arg Cys Pro 324 1 327 <210> SEQ ID NO: 10 328 <211> LENGTH: 4 329 <212> TYPE: PRT 330 <213> ORGANISM: Artificial Sequence

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Input Set : A:\iu-0008.ST25.txt

Output Set: N:\CRF4\11262002\J067122B.raw

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332 <220> FEATURE:
     333 <223> OTHER INFORMATION: Synthetic
     335 <220> FEATURE:
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     337 <222> LOCATION: (2)..(2)
     338 <223> OTHER INFORMATION: X=any amino acid
     341 <400> SEQUENCE: 10
W--> 343 Cys Xaa Cys Pro
     344 1
     347 <210> SEQ ID NO: 11
     348 <211> LENGTH: 25
     349 <212> TYPE: DNA
     350 <213> ORGANISM: Artificial Sequence
     352 <220> FEATURE:
     353 <223> OTHER INFORMATION: Synthetic
     355 <400> SEQUENCE: 11
     356 acctcgaggt cctgtgcatg tgaca
                                                                                 25
     359 <210> SEQ ID NO: 12
     360 <211> LENGTH: 25
     361 <212> TYPE: DNA
     362 <213> ORGANISM: Artificial Sequence
     364 <220> FEATURE:
     365 <223> OTHER INFORMATION: Synthetic
     367 <400> SEQUENCE: 12
     368 atgaattctt actgcaggag tgccc
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     372 <211> LENGTH: 11
     373 <212> TYPE: PRT
     374 <213> ORGANISM: Mus musculus
     376 <400> SEQUENCE: 13
     378 Cys Arg Pro Gly Gln Glu Leu Thr Lys Gln Gly
     379 1
                         5
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 11/26/2002 PATENT APPLICATION: US/10/067,122B TIME: 13:56:39

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 1253,1254,1255

Seq#:6; Xaa Pos. 5

Seq#:8; Xaa Pos. 2,3,5,6,7,8,9,10,11,12,13,15,16,17,19,20,21,23

Seq#:10; Xaa Pos. 2

VERIFICATION SUMMARY

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Input Set : A:\iu-0008.ST25.txt

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L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1200 L:247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0 L:305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0 L:309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:16 L:343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0